

## SEQUENCE LISTING

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<120> CRYSTAL OF BACTERIA CORE RNA POLYMERASE WITH RIFAMPICIN

<130> 2555-1-001

<140> UNASSIGNED  
 <141> 2001-03-09

<160> 4

<170> PatentIn version 3.0

<210> 1  
 <211> 1525  
 <212> PRT  
 <213> *Thermus aquaticus*

<220>  
 <221> X  
 <222> (1247)..(1247)  
 <223> Any amino acid can be placed at this position.

<400> 1

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Lys	Ile	Arg	Ser	Trp	Ser	Tyr	Gly	Glu	Val	Glu	Lys	Pro	Glu	Thr	Ile
	20				25					30					

Asn	Tyr	Arg	Thr	Leu	Lys	Pro	Glu	Arg	Asp	Gly	Leu	Phe	Asp	Glu	Arg
		35			40				45						

Ile	Phe	Gly	Pro	Ile	Lys	Asp	Tyr	Glu	Cys	Ala	Cys	Gly	Lys	Tyr	Lys
	50				55				60						

Arg	Gln	Arg	Phe	Glu	Gly	Lys	Val	Cys	Glu	Arg	Cys	Gly	Val	Glu	Val
65				70				75					80		

Thr	Arg	Ser	Ile	Val	Arg	Arg	Tyr	Arg	Met	Gly	His	Ile	Glu	Leu	Ala
			85				90						95		

Thr	Pro	Ala	Ala	His	Ile	Trp	Phe	Val	Lys	Asp	Val	Pro	Ser	Lys	Ile
	100					105				110					

Gly	Thr	Leu	Leu	Asp	Leu	Phe	Ala	Thr	Glu	Leu	Glu	Gln	Val	Leu	Tyr
	115				120				125						

Phe	Asn	Lys	Tyr	Ile	Val	Leu	Asp	Pro	Lys	Gly	Ala	Val	Leu	Asp	Gly
	130				135				140						

Val	Pro	Val	Glu	Lys	Arg	Gln	Leu	Leu	Thr	Asp	Glu	Glu	Tyr	Arg	Glu
145				150				155				160			

Leu	Arg	Tyr	Gly	Lys	Gln	Glu	Thr	Tyr	Pro	Leu	Pro	Ala	Gly	Val	Asp
	165					170						175			

Ala	Leu	Val	Lys	Asp	Gly	Glu	Glu	Val	Val	Lys	Gly	Gln	Glu	Leu	Ala
		180				185				190					

Pro Gly Val Val Ser Arg Met Asp Gly Val Gly Ser Leu Pro Leu Pro

195

200

205

Arg Arg Val Arg Val Asp Tyr Leu Arg Lys Glu Arg Ala Ala Leu Arg  
 210 215 220

Ile Pro Leu Ser Ala Trp Val Glu Lys Glu Pro Tyr Arg Pro Gly Glu  
 225 230 235 240

Val Leu Ala Glu Leu Ser Glu Pro Tyr Leu Phe Arg Ala Glu Glu Ser  
 245 250 255

Gly Val Val Glu Leu Lys Asp Leu Ala Glu Gly His Leu Ile Tyr Leu  
 260 265 270

Arg Gln Glu Glu Glu Val Val Ala Arg Tyr Phe Leu Pro Ala Gly Met  
 275 280 285

Thr Pro Leu Val Val Glu Gly Glu Ile Val Glu Val Gly Gln Pro Leu  
 290 295 300

Ala Glu Gly Lys Gly Leu Leu Arg Leu Pro Arg His Met Thr Ala Lys  
 305 310 315 320

Glu Val Glu Ala Glu Glu Gly Asp Ser Val His Leu Thr Leu Phe  
 325 330 335

Leu Glu Trp Thr Glu Pro Lys Asp Tyr Lys Val Ala Pro His Met Asn  
 340 345 350

Val Ile Val Pro Glu Gly Ala Lys Val Gln Ala Gly Glu Lys Ile Val  
 355 360 365

Ala Ala Ile Asp Pro Glu Glu Val Ile Ala Gln Ala Glu Gly Val  
 370 375 380

Val His Leu His Glu Pro Ala Ser Ile Leu Val Val Lys Ala Arg Val  
 385 390 395 400

Tyr Pro Phe Glu Asp Asp Val Glu Val Thr Thr Gly Asp Arg Val Ala  
 405 410 415

Pro Gly Asp Val Leu Ala Asp Gly Lys Val Lys Ser Glu Ile Tyr  
 420 425 430

Gly Arg Val Glu Val Asp Leu Val Arg Asn Val Val Arg Val Val Glu  
 435 440 445

Ser Tyr Asp Ile Asp Ala Arg Met Gly Ala Glu Ala Ile Gln Glu Leu  
 450 455 460

Leu Lys Glu Leu Asp Leu Glu Lys Leu Glu Arg Glu Leu Leu Glu Glu  
 465 470 475 480

Met Lys His Pro Ser Arg Ala Arg Arg Ala Lys Ala Arg Lys Arg Leu  
 485 490 495

Glu Val Val Arg Ala Phe Leu Asp Ser Gly Asn Arg Pro Glu Trp Met  
 500 505 510

Ile Leu Glu Ala Val Pro Val Leu Pro Pro Asp Leu Arg Pro Met Val  
 515 520 525

Gln Val Asp Gly Gly Arg Phe Ala Thr Ser Asp Leu Asn Asp Leu Tyr  
 530 535 540

Arg Arg Leu Ile Asn Arg Asn Asn Arg Leu Lys Lys Leu Leu Ala Gln  
 545 550 555 560

Gly Ala Pro Glu Ile Ile Arg Asn Glu Lys Arg Met Leu Gln Glu

565

570

575

Ala Val Asp Ala Val Ile Asp Asn Gly Arg Arg Gly Ser Pro Val Thr  
 580 585 590

Asn Pro Gly Ser Glu Arg Pro Leu Arg Ser Leu Thr Asp Ile Leu Ser  
 595 600 605

Gly Lys Gln Gly Arg Phe Arg Gln Asn Leu Leu Gly Lys Arg Val Asp  
 610 615 620

Tyr Ser Gly Arg Ser Val Ile Val Val Gly Pro Gln Leu Lys Leu His  
 625 630 635 640

Gln Cys Gly Leu Pro Lys Arg Met Ala Leu Glu Leu Phe Lys Pro Phe  
 645 650 655

Leu Leu Lys Lys Met Glu Glu Lys Ala Phe Ala Pro Asn Val Lys Ala  
 660 665 670

Ala Arg Arg Met Leu Glu Arg Gln Arg Asp Ile Lys Asp Glu Val Trp  
 675 680 685

Asp Ala Leu Glu Glu Val Ile His Gly Lys Val Val Leu Leu Asn Arg  
 690 695 700

Ala Pro Thr Leu His Arg Leu Gly Ile Gln Ala Phe Gln Pro Val Leu  
 705 710 715 720

Val Glu Gly Gln Ser Ile Gln Leu His Pro Leu Val Cys Glu Ala Phe  
 725 730 735

Asn Ala Asp Phe Asp Gly Asp Gln Met Ala Val His Val Pro Leu Ser  
 740 745 750

Ser Phe Ala Gln Ala Glu Ala Arg Ile Gln Met Leu Ser Ala His Asn  
 755 760 765

Leu Leu Ser Pro Ala Ser Gly Glu Pro Leu Ala Lys Pro Ser Arg Asp  
 770 775 780

Ile Ile Leu Gly Leu Tyr Tyr Ile Thr Gln Val Arg Lys Glu Lys Lys  
 785 790 795 800

Gly Ala Gly Met Ala Phe Ala Thr Pro Glu Glu Ala Leu Ala Ala Tyr  
 805 810 815

Glu Arg Gly Glu Val Ala Leu Asn Ala Pro Ile Val Val Ala Gly Arg  
 820 825 830

Glu Thr Ser Val Gly Arg Leu Lys Phe Val Phe Ala Asn Pro Asp Glu  
 835 840 845

Ala Leu Leu Ala Val Ala His Gly Leu Leu Asp Leu Gln Asp Val Val  
 850 855 860

Thr Val Arg Tyr Leu Gly Arg Arg Leu Glu Thr Asn Pro Gly Arg Ile  
 865 870 875 880

Leu Phe Ala Arg Ile Val Gly Glu Ala Val Gly Asp Glu Lys Val Ala  
 885 890 895

Gln Glu Leu Ile Gln Met Asp Val Pro Gln Glu Lys Asn Ser Leu Lys  
 900 905 910

Asp Leu Val Tyr Gln Ala Phe Leu Arg Leu Gly Met Glu Lys Thr Ala  
 915 920 925

Arg Leu Leu Asp Ala Leu Lys Tyr Tyr Gly Phe Thr Leu Ser Thr Thr

930

935

940

Ser Gly Ile Ile Thr Ile Gly Ile Asp Asp Ala Val Ile Pro Glu Glu  
 945 950 955 960

Lys Gln Arg Tyr Leu Glu Glu Ala Asp Arg Lys Leu Arg Gln Ile Glu  
 965 970 975

Gln Ala Tyr Glu Met Gly Phe Leu Thr Asp Arg Glu Arg Tyr Asp Gln  
 980 985 990

Val Ile Gln Leu Trp Thr Glu Thr Thr Glu Lys Val Thr Gln Ala Val  
 995 1000 1005

Phe Asn Asn Phe Glu Glu Asn Tyr Pro Phe Asn Pro Leu Tyr Val  
 1010 1015 1020

Met Ala Gln Ser Gly Ala Arg Gly Asn Pro Gln Gln Ile Arg Gln  
 1025 1030 1035

Leu Cys Gly Met Arg Gly Leu Met Gln Lys Pro Ser Gly Glu Thr  
 1040 1045 1050

Phe Glu Val Pro Val Arg Ser Ser Phe Arg Glu Gly Leu Thr Val  
 1055 1060 1065

Leu Glu Tyr Phe Ile Ser Ser His Gly Ala Arg Lys Gly Gly Ala  
 1070 1075 1080

Asp Thr Ala Leu Arg Thr Ala Asp Ser Gly Tyr Leu Thr Arg Lys  
 1085 1090 1095

Leu Val Asp Val Ala His Glu Ile Val Val Arg Glu Ala Asp Cys  
 1100 1105 1110

Gly Thr Thr Lys Tyr Ile Ser Val Pro Leu Phe Gln Met Asp Glu  
 1115 1120 1125

Val Thr Arg Thr Leu Arg Leu Arg Lys Arg Ser Asp Ile Glu Ser  
 1130 1135 1140

Gly Leu Tyr Gly Arg Val Leu Ala Arg Glu Val Glu Ala Leu Gly  
 1145 1150 1155

Arg Arg Leu Glu Glu Gly Arg Tyr Leu Ser Leu Glu Asp Val His  
 1160 1165 1170

Phe Leu Ile Lys Ala Ala Glu Ala Gly Glu Val Arg Glu Val Pro  
 1175 1180 1185

Val Arg Ser Pro Leu Thr Cys Gln Thr Arg Tyr Gly Val Cys Gln  
 1190 1195 1200

Lys Cys Tyr Gly Tyr Asp Leu Ser Met Ala Arg Pro Val Ser Ile  
 1205 1210 1215

Gly Glu Ala Val Gly Val Val Ala Ala Glu Ser Ile Gly Glu Pro  
 1220 1225 1230

Gly Thr Gln Leu Thr Met Arg Thr Phe His Thr Gly Gly Xaa Ala  
 1235 1240 1245

Val Gly Thr Asp Ile Thr Gln Gly Leu Pro Arg Val Ile Glu Leu  
 1250 1255 1260

Phe Glu Ala Arg Arg Pro Lys Ala Lys Ala Val Ile Ser Glu Ile  
 1265 1270 1275

Asp Gly Val Val Arg Ile Glu Glu Gly Glu Asp Arg Leu Ser Val

1280	1285	1290
Phe Val Glu Ser Glu Gly Phe	Ser Lys Glu Tyr Lys	Leu Pro Lys
1295	1300	1305
Asp Ala Arg Leu Leu Val Lys	Asp Gly Asp Tyr Val	Glu Ala Gly
1310	1315	1320
Gln Pro Leu Thr Arg Gly Ala	Ile Asp Pro His Gln	Leu Leu Glu
1325	1330	1335
Ala Lys Gly Pro Glu Ala Val	Glu Arg Tyr Leu Val	Asp Glu Ile
1340	1345	1350
Gln Lys Val Tyr Arg Ala Gln	Gly Val Lys Leu His	Asp Lys His
1355	1360	1365
Ile Glu Ile Val Val Arg Gln	Met Leu Lys Tyr Val	Glu Val Thr
1370	1375	1380
Asp Pro Gly Asp Ser Pro Leu	Leu Glu Gly Gln Val	Leu Glu Lys
1385	1390	1395
Trp Asp Val Glu Ala Leu Asn	Glu Arg Leu Ile Ala	Glu Gly Lys
1400	1405	1410
Val Pro Val Ala Trp Lys Pro	Leu Leu Met Gly Val	Thr Lys Ser
1415	1420	1425
Ala Leu Ser Thr Lys Ser Trp	Leu Ser Ala Ala Ser	Phe Gln Asn
1430	1435	1440
Thr Thr His Val Leu Thr Glu	Ala Ala Ile Ala Gly	Lys Lys Asp
1445	1450	1455
Glu Leu Ile Gly Leu Lys Glu	Asn Val Ile Leu Gly	Arg Leu Ile
1460	1465	1470
Pro Ala Gly Thr Gly Ser Asp	Phe Val Arg Phe Thr	Gln Val Val
1475	1480	1485
Asp Gln Arg Thr Leu Lys Ala	Ile Glu Glu Ala Arg	Lys Glu Ala
1490	1495	1500
Val Glu Ala Lys Glu Lys Glu	Ala Pro Arg Arg Pro	Val Arg Arg
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Glu Gln Pro Gly Lys Gly Leu		
1520	1525	

<210> 2  
 <211> 1119  
 <212> PRT  
 <213> *Thermus aquaticus*

<220>  
 <221> X  
 <222> (695)..(696)  
 <223> Any amino acid can be at either position.

<400> 2

Met Lys Ile Lys Arg Phe Gly Arg Ile Arg Glu Val Ile Pro Leu Pro  
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Pro Leu Thr Glu Ile Gln Val Glu Ser Tyr Lys Lys Ala Leu Gln Ala  
 20 25 30

Asp Val Pro Pro Glu Lys Arg Glu Asn Val Gly Ile Gln Ala Ala Phe  
 35 40 45  
 Lys Glu Thr Phe Pro Ile Glu Glu Gly Asp Lys Gly Lys Gly Gly Leu  
 50 55 60  
 Val Leu Asp Phe Leu Glu Tyr Arg Ile Gly Asp Pro Pro Phe Ser Gln  
 65 70 75 80  
 Asp Glu Cys Arg Glu Lys Asp Leu Thr Tyr Gln Ala Pro Leu Tyr Ala  
 85 90 95  
 Arg Leu Gln Leu Ile His Lys Asp Thr Gly Leu Ile Lys Glu Asp Glu  
 100 105 110  
 Val Phe Leu Gly His Leu Pro Leu Met Thr Glu Asp Gly Ser Phe Ile  
 115 120 125  
 Ile Asn Gly Ala Asp Arg Val Ile Val Ser Gln Ile His Arg Ser Pro  
 130 135 140  
 Gly Val Tyr Phe Thr Pro Asp Pro Ala Arg Pro Gly Arg Tyr Ile Ala  
 145 150 155 160  
 Ser Ile Ile Pro Leu Pro Lys Arg Gly Pro Trp Ile Asp Leu Glu Val  
 165 170 175  
 Glu Ala Ser Gly Val Val Thr Met Lys Val Asn Lys Arg Lys Phe Pro  
 180 185 190  
 Leu Val Leu Leu Leu Arg Val Leu Gly Tyr Asp Gln Glu Thr Leu Val  
 195 200 205  
 Arg Glu Leu Ser Ala Tyr Gly Asp Leu Val Gln Gly Leu Leu Asp Glu  
 210 215 220  
 Ala Val Leu Ala Met Arg Pro Glu Glu Ala Met Val Arg Leu Phe Thr  
 225 230 235 240  
 Leu Leu Arg Pro Gly Asp Pro Pro Lys Lys Asp Lys Ala Leu Ala Tyr  
 245 250 255  
 Leu Phe Gly Leu Leu Ala Asp Pro Lys Arg Tyr Asp Leu Gly Glu Ala  
 260 265 270  
 Gly Arg Tyr Lys Ala Glu Glu Lys Leu Gly Val Gly Leu Ser Gly Arg  
 275 280 285  
 Thr Leu Val Arg Phe Glu Asp Gly Glu Phe Lys Asp Glu Val Phe Leu  
 290 295 300  
 Pro Thr Leu Arg Tyr Leu Phe Ala Leu Thr Ala Gly Val Pro Gly His  
 305 310 315 320  
 Glu Val Asp Asp Ile Asp His Leu Gly Asn Arg Arg Ile Arg Thr Val  
 325 330 335  
 Gly Glu Leu Met Ala Asp Gln Phe Arg Val Gly Leu Ala Arg Leu Ala  
 340 345 350  
 Arg Gly Val Arg Glu Arg Met Val Met Gly Ser Pro Asp Thr Leu Thr  
 355 360 365  
 Pro Ala Lys Leu Val Asn Ser Arg Pro Leu Glu Ala Ala Leu Arg Glu  
 370 375 380  
 Phe Phe Ser Arg Ser Gln Leu Ser Gln Phe Lys Asp Glu Thr Asn Pro  
 385 390 395 400

Leu Ser Ser Leu Arg His Lys Arg Arg Ile Ser Ala Leu Gly Pro Gly  
 405 410 415  
 Gly Leu Thr Arg Glu Arg Ala Gly Phe Asp Val Arg Asp Val His Arg  
 420 425 430  
 Thr His Tyr Gly Arg Ile Cys Pro Val Glu Thr Pro Glu Gly Ala Asn  
 435 440 445  
 Ile Gly Leu Ile Thr Ser Leu Ala Ala Tyr Ala Arg Val Asp Ala Leu  
 450 455 460  
 Gly Phe Ile Arg Thr Pro Tyr Arg Arg Val Lys Asn Gly Val Val Thr  
 465 470 475 480  
 Glu Glu Val Val Tyr Met Thr Ala Ser Glu Glu Asp Arg Tyr Thr Ile  
 485 490 495  
 Ala Gln Ala Asn Thr Pro Leu Glu Gly Asp Arg Ile Ala Thr Asp Arg  
 500 505 510  
 Val Val Ala Arg Arg Gly Glu Pro Val Ile Val Ala Pro Glu Glu  
 515 520 525  
 Val Glu Phe Met Asp Val Ser Pro Lys Gln Val Phe Ser Leu Asn Thr  
 530 535 540  
 Asn Leu Ile Pro Phe Leu Glu His Asp Asp Ala Asn Arg Ala Leu Met  
 545 550 555 560  
 Gly Ser Asn Met Gln Thr Gln Ala Val Pro Leu Ile Arg Ala Gln Ala  
 565 570 575  
 Pro Val Val Met Thr Gly Leu Glu Glu Arg Val Val Arg Asp Ser Leu  
 580 585 590  
 Ala Ala Leu Tyr Ala Glu Glu Asp Gly Glu Val Val Lys Val Asp Gly  
 595 600 605  
 Thr Arg Ile Ala Val Arg Tyr Glu Asp Gly Arg Leu Val Glu His Pro  
 610 615 620  
 Leu Arg Arg Tyr Ala Arg Ser Asn Gln Gly Thr Ala Phe Asp Gln Arg  
 625 630 635 640  
 Pro Arg Val Arg Val Gly Gln Arg Val Lys Lys Gly Asp Leu Leu Ala  
 645 650 655  
 Asp Gly Pro Ala Ser Glu Glu Gly Phe Leu Ala Leu Gly Gln Asn Val  
 660 665 670  
 Leu Val Ala Ile Met Pro Phe Asp Gly Tyr Asn Phe Glu Asp Ala Ile  
 675 680 685  
 Val Ile Ser Glu Glu Leu Xaa Xaa Arg Asp Phe Tyr Thr Ser Ile His  
 690 695 700  
 Ile Glu Arg Tyr Glu Ile Glu Ala Arg Asp Thr Lys Leu Gly Pro Glu  
 705 710 715 720  
 Arg Ile Thr Arg Asp Ile Pro His Leu Ser Glu Ala Ala Leu Arg Asp  
 725 730 735  
 Leu Asp Glu Glu Gly Ile Val Arg Ile Gly Ala Glu Val Lys Pro Gly  
 740 745 750  
 Asp Ile Leu Val Gly Arg Thr Ser Phe Lys Gly Glu Gln Glu Pro Ser  
 755 760 765

Pro Glu Glu Arg Leu Leu Arg Ser Ile Phe Gly Glu Lys Ala Arg Asp  
 770 775 780  
 Val Lys Asp Thr Ser Leu Arg Val Pro Pro Gly Glu Gly Gly Ile Val  
 785 790 795 800  
 Val Gly Arg Leu Arg Leu Arg Arg Gly Asp Pro Gly Val Glu Leu Lys  
 805 810 815  
 Pro Gly Val Arg Glu Val Val Arg Val Phe Val Ala Gln Lys Arg Lys  
 820 825 830  
 Leu Gln Val Gly Asp Lys Leu Ala Asn Arg His Gly Asn Lys Gly Val  
 835 840 845  
 Val Ala Lys Ile Leu Pro Val Glu Asp Met Pro His Leu Pro Asp Gly  
 850 855 860  
 Thr Pro Val Asp Val Ile Leu Asn Pro Leu Gly Val Pro Ser Arg Met  
 865 870 875 880  
 Asn Leu Gly Gln Ile Leu Glu Thr His Leu Gly Leu Ala Gly Tyr Phe  
 885 890 895  
 Leu Gly Gln Arg Tyr Ile Ser Pro Val Phe Asp Gly Ala Thr Glu Pro  
 900 905 910  
 Glu Ile Lys Glu Leu Leu Ala Glu Ala Phe Asn Leu Tyr Phe Gly Lys  
 915 920 925  
 Arg Gln Gly Glu Gly Phe Gly Val Asp Lys Arg Glu Lys Glu Val Leu  
 930 935 940  
 Ala Arg Ala Glu Lys Leu Gly Leu Val Ser Pro Gly Lys Ser Pro Glu  
 945 950 955 960  
 Glu Gln Leu Lys Glu Leu Phe Asp Leu Gly Lys Val Val Leu Tyr Asp  
 965 970 975  
 Gly Arg Thr Gly Glu Pro Phe Glu Gly Pro Ile Val Val Gly Gln Met  
 980 985 990  
 Phe Ile Met Lys Leu Tyr His Met Val Glu Asp Lys Met His Ala Arg  
 995 1000 1005  
 Ser Thr Gly Pro Tyr Ser Leu Ile Thr Gln Gln Pro Leu Gly Gly  
 1010 1015 1020  
 Lys Ala Gln Phe Gly Gly Gln Arg Phe Gly Glu Met Glu Val Trp  
 1025 1030 1035  
 Ala Leu Glu Ala Tyr Gly Ala Ala His Thr Leu Gln Glu Met Leu  
 1040 1045 1050  
 Thr Ile Lys Ser Asp Asp Ile Glu Gly Arg Asn Ala Ala Tyr Gln  
 1055 1060 1065  
 Ala Ile Ile Lys Gly Glu Asp Val Pro Glu Pro Ser Val Pro Glu  
 1070 1075 1080  
 Ser Phe Arg Val Leu Val Lys Glu Leu Gln Ala Leu Ala Leu Asp  
 1085 1090 1095  
 Val Gln Thr Leu Asp Glu Lys Asp Asn Pro Val Asp Ile Phe Glu  
 1100 1105 1110  
 Gly Leu Ala Ser Lys Arg  
 1115

<210> 3  
 <211> 313  
 <212> PRT  
 <213> *Thermus aquaticus*

<400> 3

Met Leu Glu Ser Lys Leu Lys Ala Pro Val Phe Thr Ala Thr Thr Gln  
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Gly Asp His Tyr Gly Glu Phe Val Leu Glu Pro Leu Glu Arg Gly Phe  
 20 25 30

Gly Val Thr Leu Gly Asn Pro Leu Arg Arg Ile Leu Leu Ser Ser Ile  
 35 40 45

Pro Gly Thr Ala Val Thr Ser Val Tyr Ile Glu Asp Val Leu His Glu  
 50 55 60

Phe Ser Thr Ile Pro Gly Val Lys Glu Asp Val Val Glu Ile Ile Leu  
 65 70 75 80

Asn Leu Lys Glu Leu Val Val Arg Phe Leu Asp Pro Arg Trp Arg Thr  
 85 90 95

Thr Leu Ile Leu Arg Ala Glu Gly Pro Lys Glu Val Arg Ala Val Asp  
 100 105 110

Phe Thr Pro Ser Ala Asp Val Glu Ile Met Asn Pro Asp Leu His Ile  
 115 120 125

Ala Thr Leu Glu Glu Gly Gly Lys Leu Tyr Met Glu Val Arg Val Asp  
 130 135 140

Arg Gly Val Gly Tyr Val Pro Ala Glu Arg His Gly Ile Lys Asp Arg  
 145 150 155 160

Ile Asn Ala Ile Pro Val Asp Ala Ile Phe Ser Pro Val Arg Arg Val  
 165 170 175

Ala Phe Gln Val Glu Asp Thr Arg Leu Gly Gln Arg Thr Asp Leu Asp  
 180 185 190

Lys Leu Thr Leu Arg Ile Trp Thr Asp Gly Ser Val Thr Pro Leu Glu  
 195 200 205

Ala Leu Asn Gln Ala Val Ala Ile Leu Lys Glu His Leu Asn Tyr Phe  
 210 215 220

Ala Asn Pro Glu Ala Ser Leu Leu Pro Thr Pro Glu Val Ser Lys Gly  
 225 230 235 240

Glu Lys Arg Glu Ser Ala Glu Glu Asp Leu Asp Leu Pro Leu Glu Glu  
 245 250 255

Leu Gly Leu Ser Thr Arg Val Leu His Ser Leu Lys Glu Glu Gly Ile  
 260 265 270

Glu Ser Val Arg Ala Leu Leu Ala Leu Asn Leu Lys Asp Leu Arg Asn  
 275 280 285

Ile Pro Gly Ile Gly Glu Arg Ser Leu Glu Glu Ile Arg Gln Ala Leu  
 290 295 300

Ala Lys Lys Gly Phe Thr Leu Lys Glu  
 305 310

<210> 4  
 <211> 99

<212> PRT  
<213> *Thermus aquaticus*

<400> 4

Met Ala Glu Pro Gly Ile Asp Lys Leu Phe Gly Met Val Asp Ser Lys  
1 5 10 15

Tyr Arg Leu Thr Val Val Val Ala Lys Arg Ala Gln Gln Leu Leu Arg  
20 25 30

His Arg Phe Lys Asn Thr Val Leu Glu Pro Glu Glu Arg Pro Lys Met  
35 40 45

Arg Thr Leu Glu Gly Leu Tyr Asp Asp Pro Asn Ala Val Thr Trp Ala  
50 55 60

Met Lys Glu Leu Leu Thr Gly Arg Leu Phe Phe Gly Glu Asn Leu Val  
65 70 75 80

Pro Glu Asp Arg Leu Gln Lys Glu Met Glu Arg Leu Tyr Pro Thr Glu  
85 90 95

Glu Glu Ala